



Supplemental Figure 3.

Supplemental Figure 3. CyTOF analysis of infiltrating immune populations reveals increased frequency of Arg1⁺ myeloid subsets in the tumors of FoxP3⁺ cell-depleted mice. (A) Principal component analysis (PCA) plot and (B) Volcano plot illustrating RNA sequencing analysis of myeloid cell samples sorted from KC (cont) and KC;Foxp3^{DTR} (Treg) pancreata received 3-week DT treatment following 8 weeks post pancreatitis induction. (C) The scaled heatmap showing relative marker expression in each cell population identified in viSNE map. (D) Frequencies of SPADE-identified subpopulations between groups were compared with two-tailed t-test and corrected for multiple comparisons with Benjamini-Hochberg correction with FDR = 0.1. The statistically significant populations after correction are labeled in red, n = 3 mice/group, data is displayed as mean ± SD.